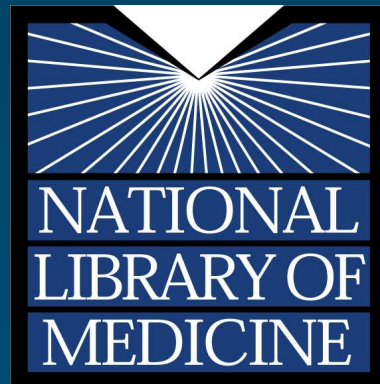


7<sup>th</sup> International Protégé Conference  
2<sup>nd</sup> Workshop on Visualizing Information  
in Knowledge Engineering (VIKE'04)

Bethesda, MD

July 6, 2004

# Biomedical Knowledge Visualization



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for Biomedical Communications  
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# UMLS Semantic Navigator

## *SemNav*

<http://umlsks.nlm.nih.gov>\*

► SN Resources ► Semantic Navigator  
(\* free UMLS registration required)

# Unified Medical Language System®

- ◆ Developed at NLM since 1990
- ◆ 15<sup>th</sup> edition in 2004
- ◆ Integrates some 60 terminological resources
  - Clinical vocabularies (including specialties)
  - Core terminologies (anatomy, drugs, med. devices)
  - Administrative terminologies, standards
- ◆ Integration
  - Synonymous terms are clustered in a concept
  - Hierarchies (trees) are combined in a graph structure



# Terminology integration Terms

Duchenne muscular dystrophy

MeSH, SNOMED  
CTV3, Jablonski,  
CRISP, DxPlain,  
MedDRA, LOINC

Duchenne's muscular dystrophy

COSTAR

Duchenne de Boulogne muscular dystrophy

Jablonski

Duchenne type progressive muscular dystrophy

SNOMED

pseudohypertrophic muscular dystrophy

MeSH, CTV3  
SNOMED

X-linked recessive muscular dystrophy

Jablonski

severe generalized familial muscular dystrophy

SNOMED

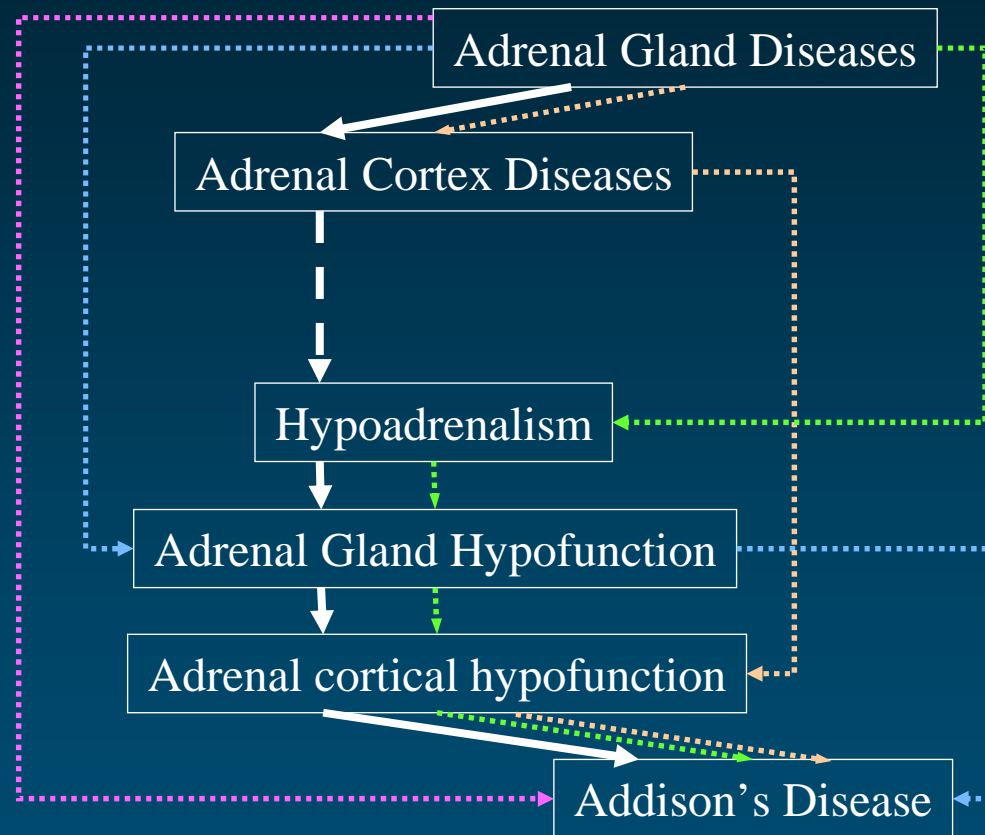


# Terminology integration

## Relationships

SNOMED  
MeSH  
AOD  
Read Codes

UMLS



# UMLS

## ◆ Two-level structure

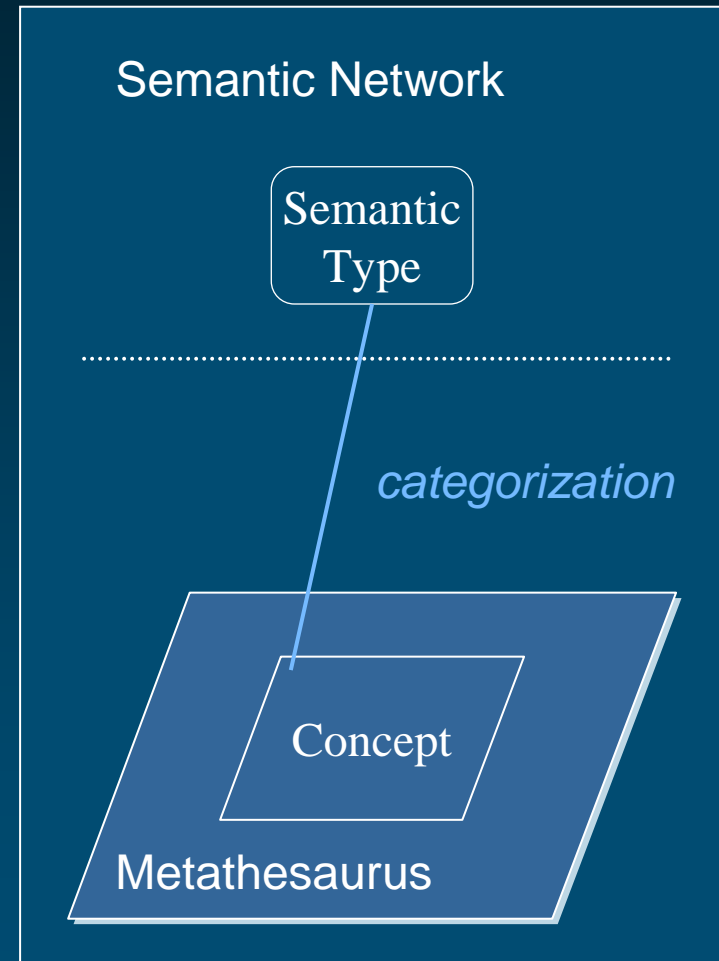
### ● Semantic Network

- 135 Semantic Types (STs)
- 54 types of relationships among STs

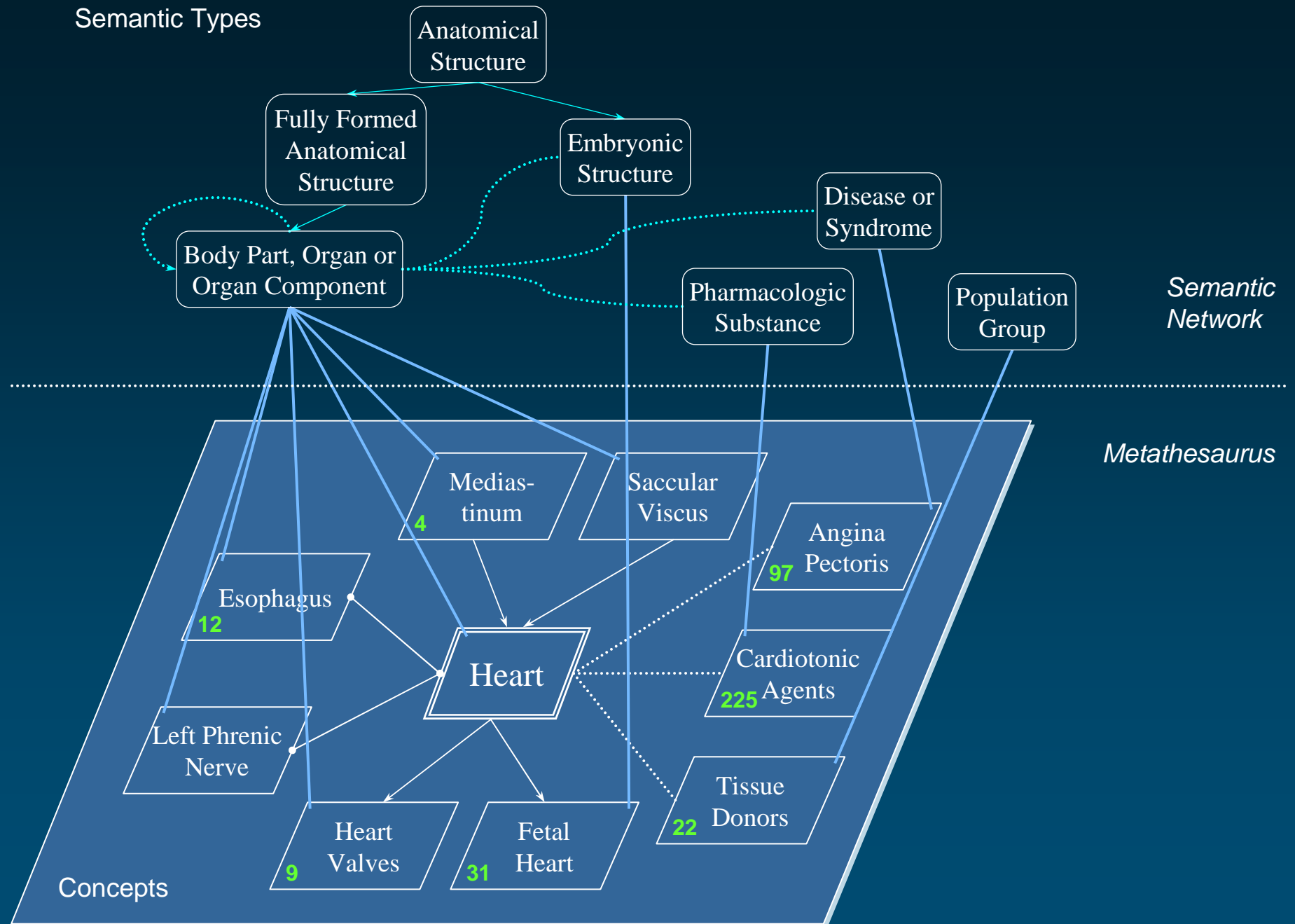
### ● Metathesaurus

- >1M concepts
- ~12 M inter-concept relationships

### ● Link = categorization



## Semantic Types









[Amino Acids, Peptides, and Proteins \[D12\]](#)

[Proteins \[D12.776\]](#)

[Contractile Proteins \[D12.776.210\]](#)

[Muscle Proteins \[D12.776.210.500\]](#)

[Actinin \[D12.776.210.500.095\]](#)

[Actins \[D12.776.210.500.100\]](#)

[Actomyosin \[D12.776.210.500.154\]](#)

[Calsequestrin \[D12.776.210.500.220\]](#)

► [Dystrophin \[D12.776.210.500.250\]](#)

[Myogenic Regulatory Factors \[D12.776.210.500.570\] +](#)

[Myoglobin \[D12.776.210.500.588\]](#)

[Myosins \[D12.776.210.500.600\] +](#)

[Parvalbumins \[D12.776.210.500.750\]](#)

[Ryanodine Receptor Calcium Release Channel \[D12.776.210.500.800\]](#)

[Tropomyosin \[D12.776.210.500.895\]](#)

[Troponin \[D12.776.210.500.910\] +](#)

[Amino Acids, Peptides, and Proteins \[D12\]](#)

[Proteins \[D12.776\]](#)

[Cytoskeletal Proteins \[D12.776.220\]](#)

[Adenomatous Polyposis Coli Protein \[D12.776.220.040\]](#)

► [Dystrophin \[D12.776.220.250\]](#)

[Intermediate Filament Proteins \[D12.776.220.475\] +](#)

[Microfilament Proteins \[D12.776.220.525\] +](#)

[Microtubule Proteins \[D12.776.220.600\] +](#)

[Spectrin \[D12.776.220.980\]](#)

[Talin \[D12.776.220.985\]](#)

[Vinculin \[D12.776.220.990\]](#)

# MeSH Browser



[Amino Acids, Peptides, and Proteins \[D12\]](#)

[Proteins \[D12.776\]](#)

[Membrane Proteins \[D12.776.543\]](#)

[Ankyrins \[D12.776.543.080\]](#)

[Arrestins \[D12.776.543.090\] +](#)

[Bacterial Outer Membrane Proteins \[D12.776.543.100\] +](#)

[Caveolins \[D12.776.543.160\]](#)

[Clathrin \[D12.776.543.200\]](#)

[Coat Protein Complex I \[D12.776.543.212\] +](#)

[Connexins \[D12.776.543.225\] +](#)

► [Dystrophin \[D12.776.543.250\]](#)

[Heterotrimeric GTP-Binding Proteins \[D12.776.543.325\] +](#)

[LDL-Receptor Related Protein-Associated Protein \[D12.776.543.475\]](#)

[Membrane Glycoproteins \[D12.776.543.550\] +](#)

[Membrane Transport Proteins \[D12.776.543.585\] +](#)

[Myelin Proteins \[D12.776.543.620\] +](#)

[Neurofibromin 2 \[D12.776.543.685\]](#)

[Receptors, Cell Surface \[D12.776.543.750\] +](#)

[Spectrin \[D12.776.543.980\]](#)



# SemNav Visualization options

Start again    Apply new parameters

**Restrict to vocabulary:** Show all ▼

**Highlight vocabulary:** Nothing ▼

**UMLS data:** UMLS\_2002 ▼

**Type of hierarchical rel:** ☒ All ☐ Parent/Child only ☐ Broader/Narrower only

**Transitive reduction:** ☒ yes ☐ no

Start again    Apply new parameters







UMLS\_2002} UMLS® Semantic Navigator [2.07] - Netscape 6

File Edit View Search Go Bookmarks Tasks Help

http://etbsun2.nlm.nih.gov:8000/perl/semmav.cgi.pl

Search

Siblings

Chemicals & Drugs

- (LA)12 peptide
- (methyl)ammonium uptake carrier
- Corynebacterium
- 120-kDa hemocyte-specific membrane protein, flesh fly
- 15a protein, Aedes aegypti
- 22.6-kDa antigen, Schistosoma japonicum
- 36-kDa vesicular integral membrane protein
- 38L protein
- 5-lipoxygenase-activating protein
- 59 kDa dystrophin-associated protein
- A-1 antigen
- A-kinase anchor protein 149
- A-kinase anchor protein 15
- A-kinase anchor protein 200
- A-kinase anchor protein KL
- A14.5L protein
- A15 protein
- A4 protein
- ABC-me protein
- AcB protein
- ACR3 protein
- AcEt protein
- actA protein
- Actinin

Other Related Concepts

Disorders

- muscular dystrophy
- Muscular Dystrophy, Duchenne

Living Beings

- Mice, Inbred mdx
- (3 other related concepts)

Co-occurring Concepts

Anatomy

- Brain
- Cell Membrane
- Cytoskeleton
- Heart
- Hippocampus
- Muscle
- Fibers
- Muscle Fibers, Fast-Twitch
- Muscle, Skeletal
- Muscle, Smooth
- Muscle, Smooth, Vascular

apodystrophin 1

140-kDa dystrophin

dystrophin-related protein 2

dys-1 protein

Dp260 protein

proteins by body part

Microfilament Proteins

membrane protein

actin binding protein

binding protein

contractile protein

muscle protein

dystrophin

Start again

Apply new parameters

Restrict to vocabulary:

Show all

Highlight vocabulary:

CRISP

UMLS data:

UMLS\_2002

Type of hierarchical rel:

All

Parent/Child only

Broader/Narrower only

Transitive reduction:

yes

no

Similar Concepts

(none)

Closest MeSH Terms

Main Headings

- Dystrophin

Subheadings

LEGEND

Dystrophin

Document: Done (68.78 sec)



UMLS 2002J UMLS® Semantic Navigator » [2.07] - Netscape 6

File Edit View Search Go Bookmarks Tasks Help

http://atbsun2.nlm.nih.gov:8000/cgi/semnav.cgi.pl

search

Siblings

Disorders

- Erb's muscular dystrophy
- Gower's muscular dystrophy
- Intermediate X-linked muscular dystrophy
- Manifesting female carrier of X-linked muscular dystrophy
- Muscular Dystrophy, Becker
- Muscular Dystrophy, Facioscapulohumeral
- Muscular Dystrophy, Lamb-Girdle
- Muscular Dystrophy, Oculopharyngeal
- Ocular muscular dystrophy

(9 siblings)

[direct children and narrower concepts of direct parents and broader concepts]

Other Related Concepts

Chemicals & Drugs

- Dystrophin

Disorders

- Muscular Dystrophy, Becker
- MUSCULAR DYSTROPHY, PSEUDOHYPERTROPHIC, PROGRESSIVE, DUCHENNE AND BECKER TYPES
- Progressive muscular dystrophy, legs

(4 other related concepts)

Co-occurring Concepts

Activities & Behaviors

- Knowledge, Attitudes, Practice
- Physician's Practice Patterns
- Psychological adjustment
- Verbal Behavior

Anatomy

- Achilles Tendon
- Ankle

X-linked muscular dystrophy with limb girdle distribution

Muscular Dystrophies

Muscular dystrophies and other myopathies

X-linked muscular dystrophy with abnormal dystrophin

Hereditary progressive muscular dystrophy, NOS

Muscular Dystrophy, Duchenne

Muscular Dystrophy, Duchenne

Start again

Restrict to vocabulary

Highlight vocabulary

UMLS date

Type of hierarchical rel.

Transitive reduction

Apply new parameters

Show all

Nothing

UMLS\_2002

All

Parent/Child only

Broader/Narrower only

yes

no

Muscular Dystrophy, Duchenne

LEGEND

Closest MeSH Terms

Main Headings

- Muscular Dystrophy, Duchenne

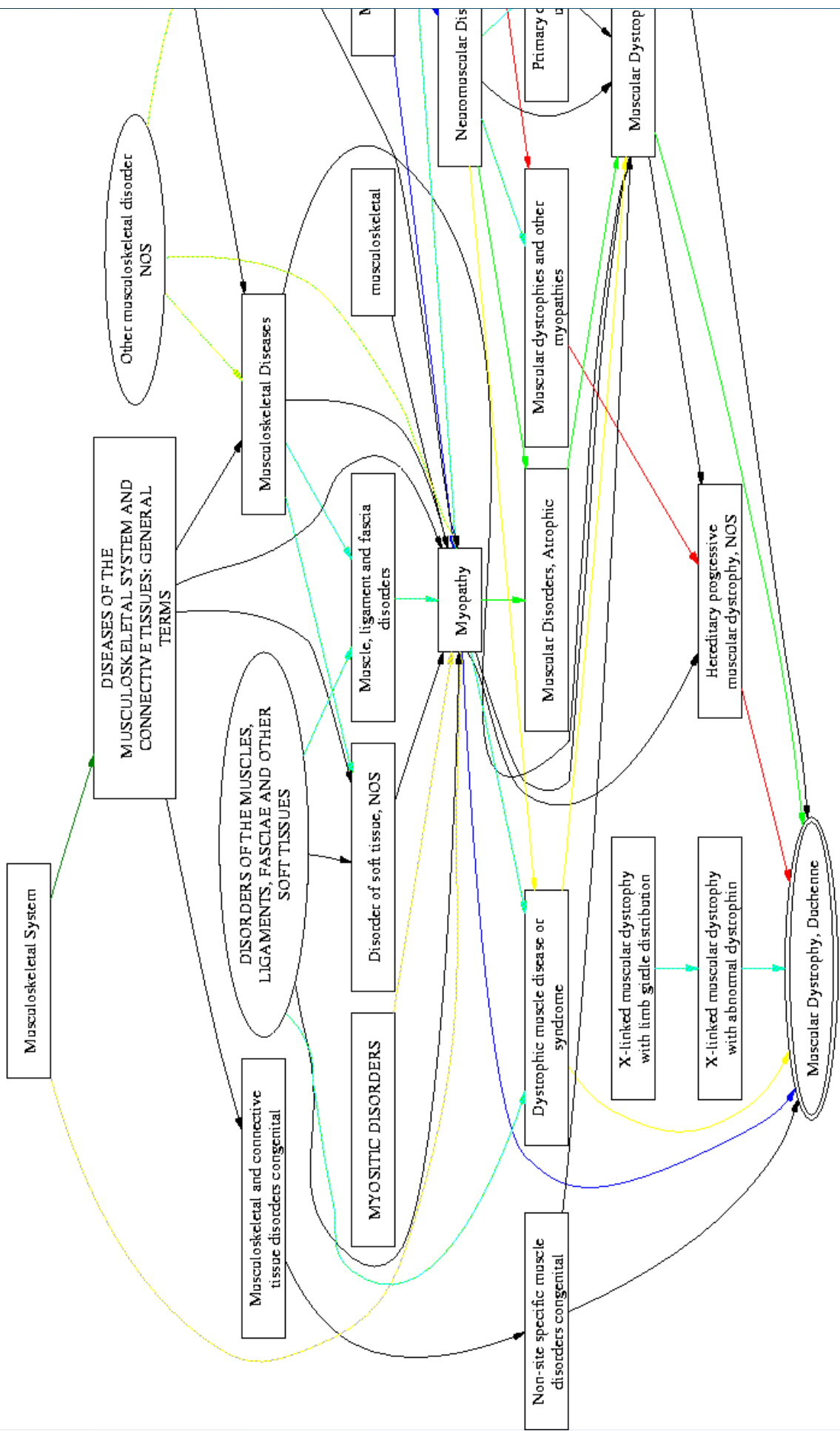
Subheadings

Similar Concepts

Hereditary progressive muscular dystrophy, NOS

(1 similar concept)

Document: Done (12,406 secs)





# SemNav R

Relationships of **Dystrophin** (C1)  
*Amino Acid, Peptide, or Protein*  
*Biologically Active Substance*  
 to **Muscular Dystrophy, Duchenne** (C2)  
*Disease or Syndrome*

---

**Metathesaurus Relationships**

C1 *otherwise related to* C2  
 • MeSH

C1 *co-occurs with* C2  
 Frequency = 55 • MEDLINE

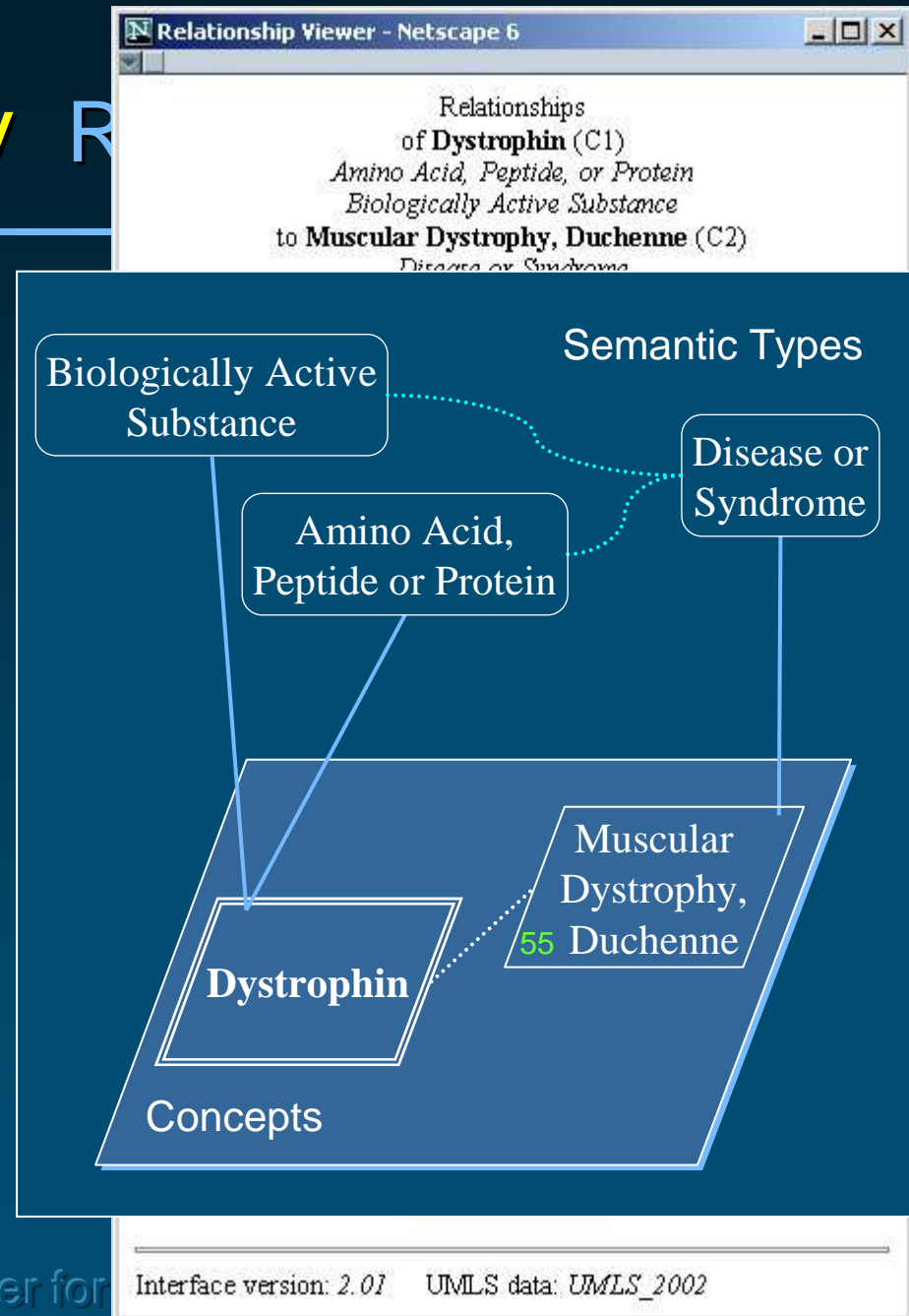
---

**Semantic Network Relationships**

Amino Acid, Peptide, or Protein	<ul style="list-style-type: none"> <li>affects</li> <li>causes</li> </ul>	Disease or Syndrome
Biologically Active Substance	<ul style="list-style-type: none"> <li>affects</li> <li>causes</li> <li>complicates</li> <li>produced_by</li> </ul>	Disease or Syndrome

[Close this window](#)

Interface version: 2.01 UMLS data: UMLS\_2002



# Gene Ontology browser

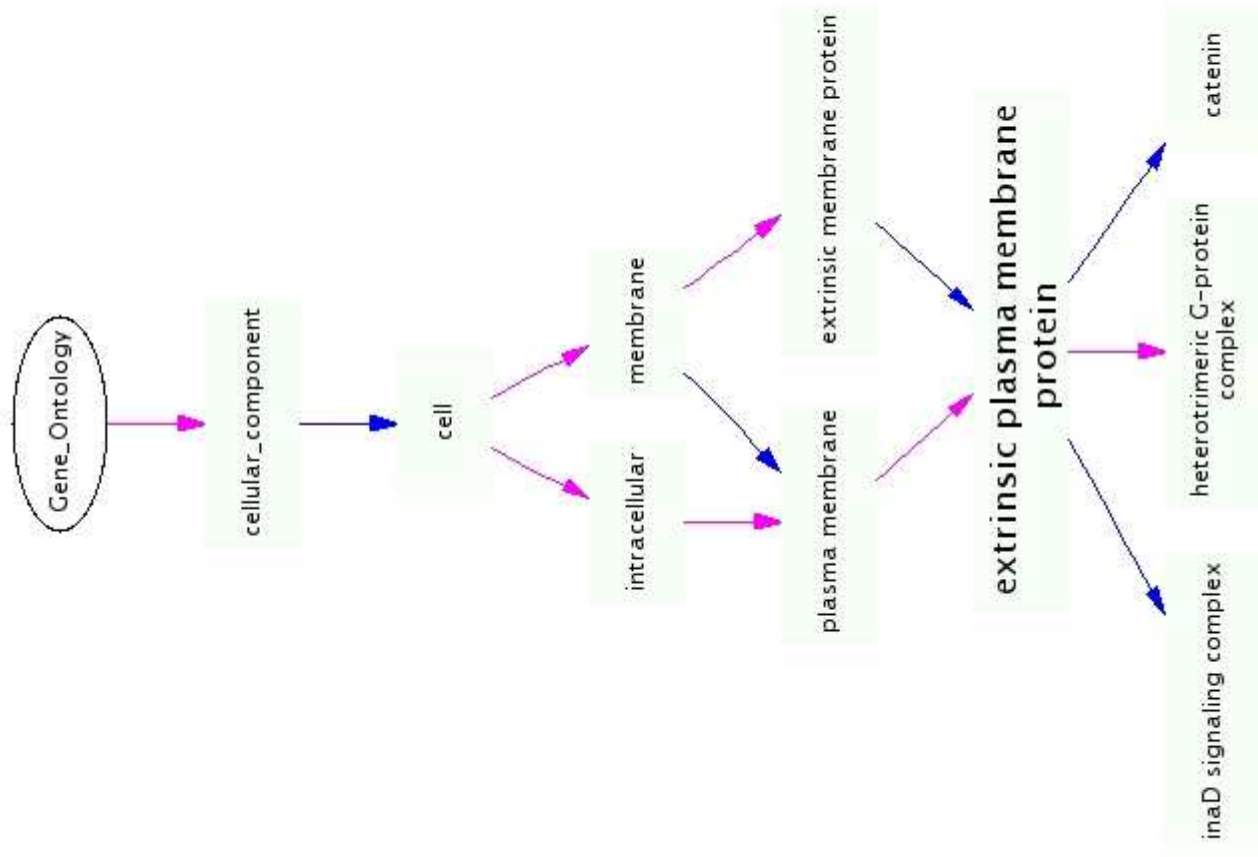


<http://mor.nlm.nih.gov/perl/gennav.pl>

# Gene Ontology™

- ◆ Developed by the GO Consortium
- ◆ Several components (GO database)
  - Ontology (~17,000 concepts)
    - Molecular functions
    - Cellular components
    - Biological processes
  - Gene products (~1.6M)
  - Associations between Gene products and GO concepts (~6.8M)





AmiGO : Tree View - Netscape 6

File Edit View Search Go Bookmarks Tasks Help

http://www.godatabase.org/cgi-bin/go.cgi?action=

Home Search Bookmarks Google NLM P Biblio Br

AmiGO

Search GO:

Top Docs [Gene Ontology GO Links](#) [GO Summary](#) [Terms](#) [Gene Products](#)

**GO:0003673 : Gene Ontology (33650)**

- + GO:0008150 : biological process (24768)
- + GO:0005575 : cellular component (17255)
- + GO:0005623 : cell (14268)
  - + GO:0005622 : intracellular (12771)
    - + GO:0005886 : plasma membrane (2273)
      - + GO:0019897 : extrinsic plasma membrane protein (4511)
        - + GO:0019898 : extrinsic membrane protein (58)
        - + GO:0019897 : extrinsic plasma membrane protein (4511)
          - + GO:0005886 : plasma membrane (2273)
            - + GO:0019897 : extrinsic plasma membrane protein (4511)
              - + GO:0003674 : molecular function (23707)

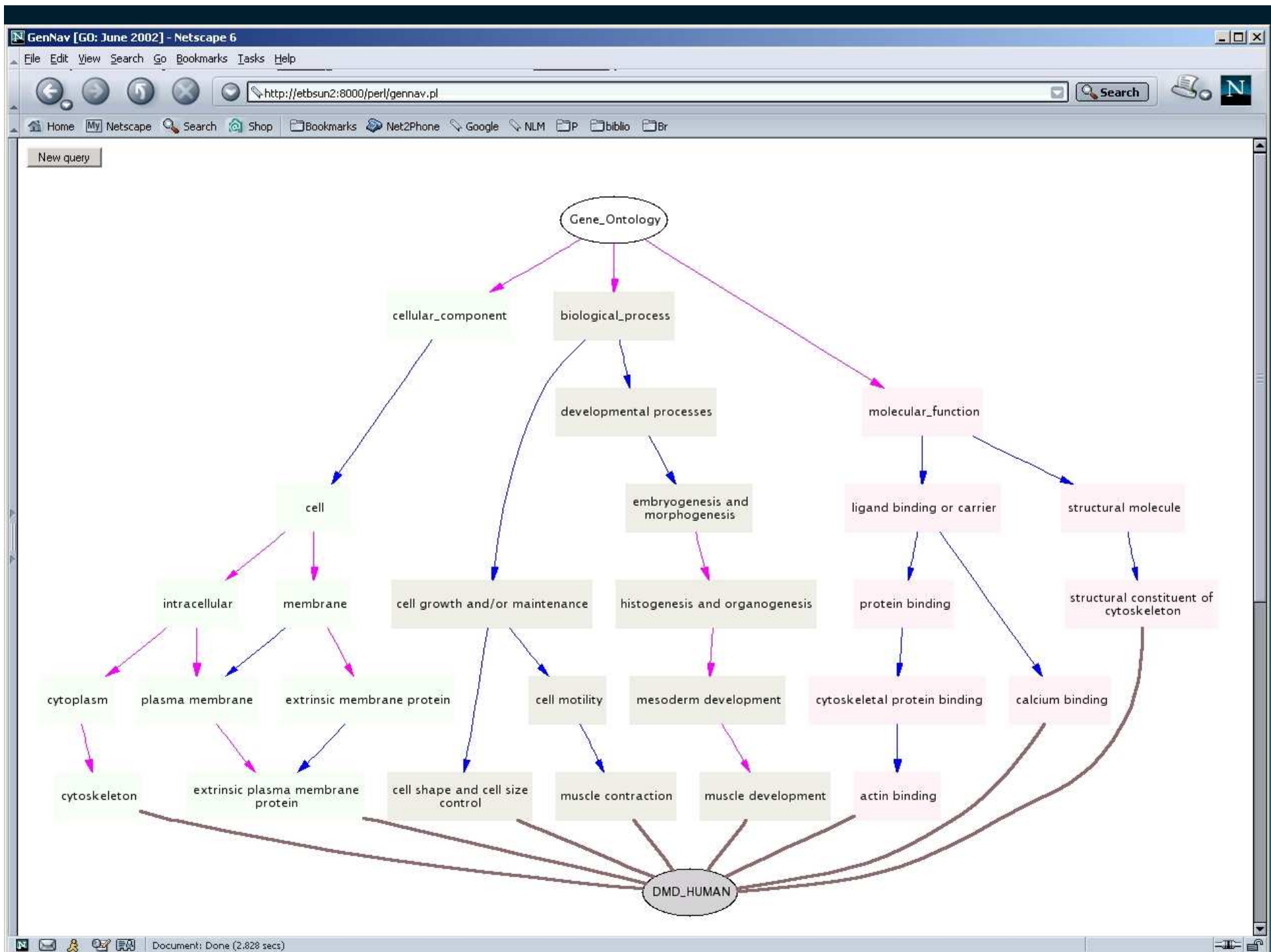
Get this GO tree as RDF XML.

Get this data as a GO flat file.

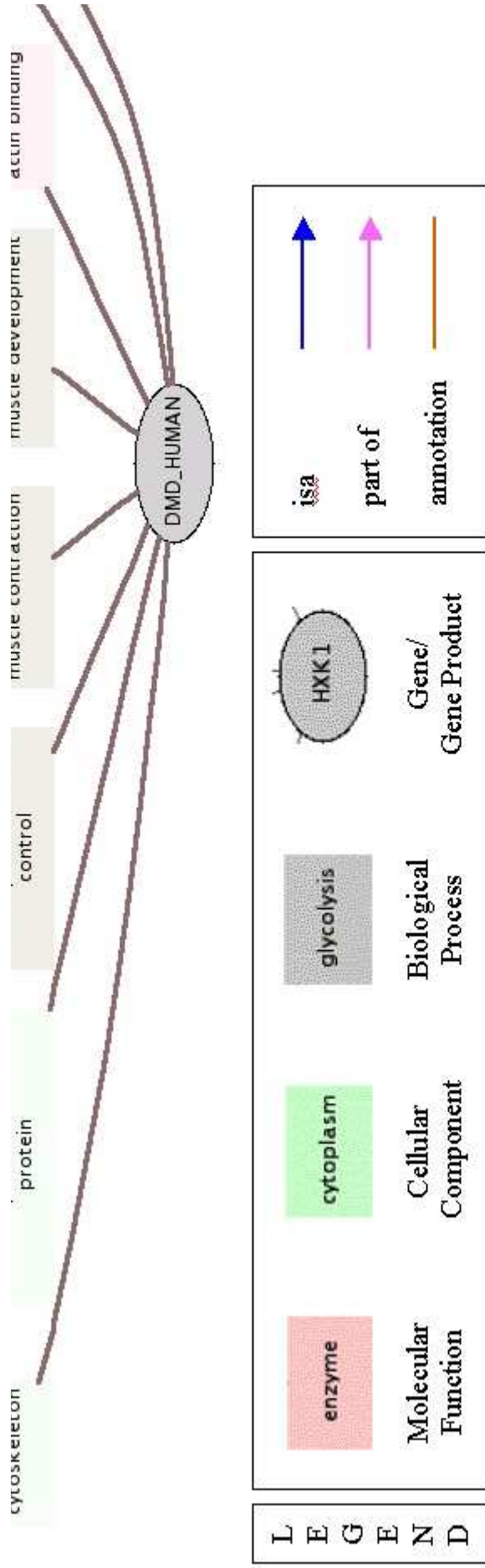
Get a bookmarkable url of this GO tree.

Copyright The Gene Ontology Consortium. All rights reserved.

Document: Done (2.437 secs)



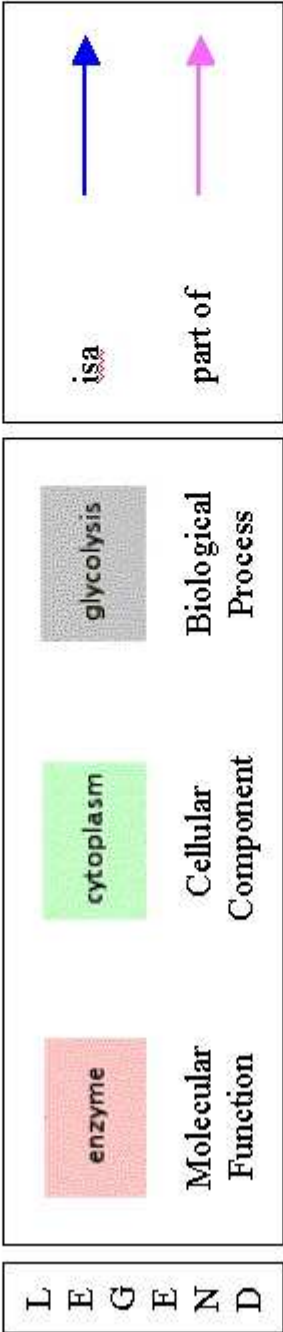
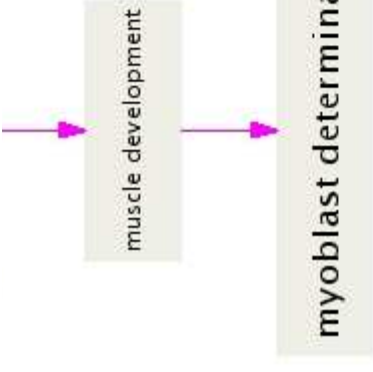




Symbol	DMD_HUMAN				
Species	0				
Full name	Dystrophin				
Synonyms	<ul style="list-style-type: none"> <li>• IPI00006091</li> </ul>				
Cross-references	DB TYPE	ID	OBJ_TYPE	TAXON	
	sp	acc	P11532	protein	9606
Molecular functions	<ul style="list-style-type: none"> <li>• actin binding [IEA]</li> <li>• structural constituent of cytoskeleton [TAS]</li> <li>• calcium binding [IEA]</li> </ul>				
Biological processes	<ul style="list-style-type: none"> <li>• muscle contraction [NR]</li> <li>• cell shape and cell size control [TAS]</li> <li>• muscle development [NR]</li> </ul>				
Cellular components	<ul style="list-style-type: none"> <li>• extrinsic plasma membrane protein [TAS]</li> <li>• cytoskeleton [TAS]</li> </ul>				

New query

Interface version: 0.3



Name	myoblast determination		
Type	process		
Accession number	7518		
Synonyms	none		
Definitions	none		
Parents (isa)	none		
Parents (part of)	• muscle development		
Children (isa)	none		
Children (part of)	none		
Cross-references	ID KEY DB		
Gene products (restricted to species: 9606)	Full name		Evidence
	Interferon-related developmental regulator 1	Symbol IFR1_HUMAN	Taxon 9606
	MEF-2 PROTEIN	Symbol Q9Y655	Taxon 9606

# GenNav

**Term:**

**What:**

**Field:**

**String matching method:**

**Species restriction:**

Search GO Clear

[Comments? Feedback? Questions?](#)

Contact: [Olivier Bodenreider](#)

[Copyright and Privacy Notice](#)

Interface version: 0.3



# Technical details

# Technical details

- ◆ Simple web/cgi technology (apache, Perl)
- ◆ dot (GraphViz)
  - PNG file (-Tpng)
  - Client-side map (-Tcmap)
- ◆ Precompute the transitive closure on hierarchical relations to perform the transitive closure fast
- ◆ Remove cycles (UMLS)



# Discussion

## *Issues and Challenges*

# Issues

## ◆ Size

- Large number of concepts (>1 million)

## ◆ Complexity

- Polyhierarchical structures
- Multiple information sources
- Multiple properties

## ◆ Lack of formality

- Redundant relations
- Hierarchies vs. hierarchical relations



# Challenges

- ◆ Restrict information space
  - To selected information sources (SemNav)
  - To selected organisms (GenNav)
- ◆ Reduce complexity (SemNav)
  - Group concepts by semantic groups
  - Transitive reduction on hierarchical relations
  - Select co-occurring concepts
- ◆ Reduce the cognitive burden on the user
  - Use graph-based rather than tree-based representations



# SemNav Semantic groups

The screenshot displays the SemNav Semantic Navigator interface. The main window shows a hierarchical diagram of Addison's Disease. The diagram is structured as follows:

- Adrenal Gland Diseases**
  - Disorder definition**
    - Syndrome**
      - Addison's disease with adrenocortical insufficiency**
      - Addison's Disease Secondary to Idiopathic Atrophy**

The diagram also includes a list of **Siblings** on the left, which includes various disorders such as Acquired Immune deficiency Syndrome, Addison's disease, and Addison's disease with adrenocortical insufficiency. The right side of the interface shows a list of **Other Related Concepts**, including Addison's disease, Addison's disease with adrenocortical insufficiency, and Addison's disease secondary to idiopathic atrophy. The bottom of the interface features a search bar and a list of **Co-occurring Concepts**, which are categorized into **Anatomy** and **Chemicals & Drugs**.

**Co-occurring Concepts**

**Anatomy**

- Adrenal Cortex [12] ☐
- Adrenal Glands [19] ☐
- Ear Cartilages [2] ☐
- Ear, External [2] ☐
- Liver [2] ☐
- Pituitary Gland [3] ☐
- Tears body substance [2] ☐
- X Chromosome [3] ☐

**Chemicals & Drugs**

- Alanine Transaminase [2] ☐
- Aldosterone [3] ☐
- Anti-Inflammatory Agents, Steroidal [2] ☐
- Antigens,

# Challenges

- ◆ Restrict information space
  - To selected information sources (SemNav)
  - To selected organisms (GenNav)
- ◆ Reduce complexity (SemNav)
  - Group concepts by semantic groups
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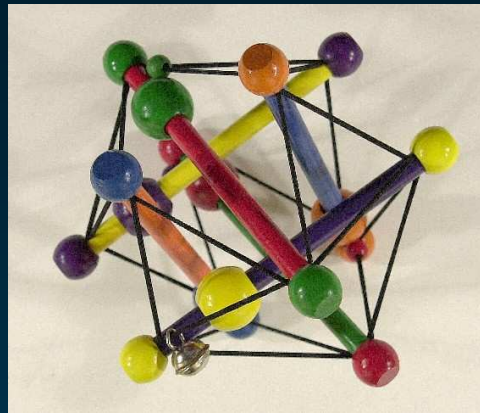




# Challenges

- ◆ Restrict information space
  - To selected information sources (SemNav)
  - To selected organisms (GenNav)
- ◆ Reduce complexity (SemNav)
  - Group concepts by semantic groups
  - Transitive reduction on hierarchical relations
  - Select co-occurring concepts
- ◆ Reduce the cognitive burden on the user
  - Use graph-based rather than tree-based representations





# Medical Ontology Research

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